



[Marked up preliminary amendment]



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- 1) (Amended) A c[C]himeric expression promoter comprising at least one nucleic acid sequence, derived from a first plant promoter comprising a plant vascular expression promoter region, said plant vascular expression promoter region being replaced with a nucleic acid sequence derived from a second plant promoter and comprising a plant green tissue expression promoter region.
- 2) (Amended) The c[C]himeric expression promoter according to claim 1, wherein said first plant promoter originates from the Commelina Yellow Mottle Virus [(CoYMV)] and said second plant promoter originates from the Cassava Vein Mosaic Virus [(CsVMV)].
- 3) (Amended) The c[C]himeric promoter [according to any one of the preceding claims] of claim 1, wherein [the] said nucleic acid sequences originate from the intergenic regions of said first and second promoters.
- 4) (Amended) The c[C]himeric promoter [according to any one of the preceding claims] of claim 1, wherein it comprises at least a part of a nucleic acid sequence with SEQ ID No.1 [identified in the sequence listing under the number SEQ.ID01] fused to at least a part of a nucleic [identified in the sequence listing under the number SEQ.ID02] acid sequence with SEQ ID No.2.
- 5) (Amended) The c[C]himeric promoter [according to anyone of the preceding claims] of claim 1, wherein the nucleic acid sequence of [the] said chimeric promoter consists of a sequence selected from the group consisting of the sequences with SEQ ID Nos. 3-7 and 19-25. [identified in the sequence listing under the numbers SEQ.ID03, SEQ.ID04, SEQ.ID05, SEQ.ID06, SEQ.ID07, SEQ.ID19, SEQ.ID20, SEQ.ID21, SEQ.ID22, SEQ.ID23, SEQ.ID24 and SEQ.ID25.]
- 6) (Amended) A c[C]himeric expression promoter comprising a promoter of viral origin, of which a part consists of an exogenous element [capable of promoting] which promotes expression in plant green tissues [(GT)].

- 7) (Amended) The c[C]himeric expression promoter [according to] of claim 6, wherein [the] said [GT] exogenous promoter element is also of viral origin.
- 8) (Amended) The c[C]himeric expression promoter [according to] of claim [6] 7, wherein [the] said promoter of viral origin originates from the Commelina Yellow Mottle Virus [(CoYMV)].
- 9) (Amended) The c[C]himeric expression promoter [according to] of claim [8] 7, wherein [the] said exogenous promoter element originates from the Cassava Vein Mosaic Virus [(CsVMV)].
- 10) (Amended) The c[C]himeric expression promoter [according to] of claim 6, wherein the exogenous [GT] element replaces an endogenous vascular tissue expression [(VT)] promoter of viral origin.
- 11) (Amended) The c[C]himeric promoter [according to any one of the preceding claims] of claim 1 or 6, wherein it further comprises at least one “endosperm like” box.
- 12) (Amended) The c[C]himeric promoter [according to any one of the preceding claims] of claim 1 or 6, wherein it further comprises at least one “as1 like” box[ed] operably linked to [the] a plant green tissue expression GT promoter element.
- 13) (Amended) The c[C]himeric promoter [according to any one of the preceding claims] of claim 1 or 6, wherein it further comprises at least one “as1” box operably linked to [the] a green tissue expression GT promoter element.
- 14) (Amended) The c[C]himeric promoter [according to any one of the preceding claims] of claim 1 or 6, wherein it further comprises at least one “as2” box operably linked to [the] a plant green tissue expression GT promoter element.
- 15) (Amended) The c[C]himeric promoter [according to any one of the preceding claims] of claim 1 or 6, wherein [the one or more of the] said “as1 like”, “as1”, and “as2” boxes are operably linked upstream or downstream of the plant green tissue expression GT promoter element.

16) (Amended) The c[C]himeric promoter [according to any one of the preceding claims] of claim 1 or 6, wherein [the one or more of the] said “as1 like”, “as1”, and “as2” boxes are operably linked in [normal (5'>3')] 5'>3' or [inverse (3'>5')] 3'>5' orientation.

17) (Amended) The c[C]himeric promoter [according to any one of the preceding claims] of claim 1 or 6, wherein it comprises at least one “as2/as2/as2” box in [normal (5'>3')] 5'>3' or [inverse (3'>5')] 3'>5' orientation.

18) (Amended) The c[C]himeric promoter [according to any one of claims 6 to 17] of claim 6, wherein it comprises at least [a] one sequence selected from the group consisting of [the] sequences with SEQ ID Nos. 3-7 and 19-25. [identified in the sequence listing under the numbers SEQ.ID03, SEQ.ID04, SEQ.ID05, SEQ.ID06, SEQ.ID07, SEQ.ID19, SEQ.ID20, SEQ.ID21, SEQ.ID22, SEQ.ID23, SEQ.ID24 and SEQ.ID25.]

19) (Amended) An e[E]xpression cassette comprising at least one nucleic acid sequence derived from a first plant promoter comprising a plant vascular expression promoter region, said plant vascular expression promoter region being replaced with a nucleic acid sequence derived from a second plant promoter and comprising a plant green tissue expression promoter region, [the sequences] said at least one sequence being operably linked to a nucleic acid sequence or gene coding for a polypeptide to be produced, said nucleic acid sequence or gene itself operably linked to a transcription termination nucleic acid sequence.

20) (Amended) The e[E]xpression cassette [according to] of claim 19, wherein said first plant promoter originates from the Commelina Yellow Mottle Virus [(CoYMV)] and said second plant promoter originates from the Cassava Vein Mosaic Virus [(CsVMV)].

21) (Amended) The e[E]xpression cassette [according to] of claim 19, wherein it comprises at least a part of a nucleic acid sequence with SEQ ID No. 1 [identified in the sequence listing under the number SEQ.ID01] fused to at least a part of a nucleic acid sequence with SEQ ID No. 2. [identified in the sequence listing under the number SEQ. ID 02.]

22) (Amended) The e[E]xpression cassette [according to] of claim 19, wherein [the] said first or second promoter [nucleic acid sequence consists of] comprises a sequence selected from the group consisting of the sequences with SEQ ID Nos. 3-7 and 19-25. [identified in the sequence

listing under the numbers SEQ.ID03, SEQ.ID04, SEQ.ID05, SEQ.ID06, SEQ.ID07, SEQ.ID19, SEQ.ID20, SEQ.ID21, SEQ.ID22, SEQ.ID23, SEQ.ID24 and SEQ.ID25.]

26) (Canceled) Directional desoxynucleotide building block [for the construction of a chimeric expression promoter or an isolated promoter nucleic acid sequence according to any one of claims 1 to 18 or 23 or 24, wherein it corresponds to the sequence identified in the sequence listing under the number SEQ. ID 09.

27) (Canceled) Directional desoxynucleotide building block for the construction of a chimeric expression promoter or an isolated promoter nucleic acid sequence according to any one of claims 1 to 18 or 23 or 24, wherein it corresponds to the sequence identified in the sequence listing under the number SEQ. ID 10.

28) (Canceled) Directional desoxynucleotide building block for the construction of a chimeric expression promoter or an isolated promoter nucleic acid sequence according to any one of claims 1 to 18 or 23 or 24, wherein it corresponds to the sequence identified in the sequence listing under the number SEQ. ID 11.

29) (Canceled) Directional desoxynucleotide building block for the construction of a chimeric expression promoter or an isolated promoter nucleic acid sequence according to any one of claims 1 to 18 or 23 or 24, wherein it corresponds to the sequence identified in the sequence listing under the number SEQ. ID 12.

30) (Canceled) Directional desoxynucleotide building block for the construction of a chimeric expression promoter or an isolated promoter nucleic acid sequence according to any one of claims 1 to 18 or 23 or 24, wherein it corresponds to the sequence identified in the sequence listing under the number SEQ. ID 13.

31) (Canceled) Directional de[s]oxynucleotide building block for the construction of a chimeric expression promoter or an isolated promoter nucleic acid sequence according to any one of claims 1 to 18 or 23 or 24, wherein it corresponds to the sequence identified in the sequence listing under the number SEQ. ID 14.

32) (Amended) A g[G]uide de[s]oxynucleotide building block for the construction of a chimeric expression promoter or an isolated promoter nucleic acid sequence according to claim 1, 6, or 49, wherein said sequence is selected from the group consisting of SEQ ID Nos. 15-18. [any one of claims 1 to 18 or 23 or 24, wherein it corresponds to the sequence identified in the sequence listing under the number SEQ. ID 15.]

33) (Canceled) Guide desoxynucleotide building block for the construction of a chimeric expression promoter or an isolated promoter nucleic acid sequence according to any one of claims 1 to 18 or 23 or 24, wherein it corresponds to the sequence identified in the sequence listing under the number SEQ. ID No. 16.

34) (Canceled) Guide desoxynucleotide building block for the construction of a chimeric expression promoter or an isolated promoter nucleic acid sequence according to any one of claims 1 to 18 or 23 or 24, wherein it corresponds to the sequence identified in the sequence listing under the number SEQ. ID 17.

35) (Canceled) Guide desoxynucleotide building block for the construction of a chimeric expression promoter or an isolated promoter nucleic acid sequence according to any one of claims 1 to 18 or 23 or 24, wherein it corresponds to the sequence identified in the sequence listing under the number SEQ. ID 18.

36) (Amended) A v[V]ector comprising a promoter, or a promoter nucleic acid sequence, which [capable of initiating] initiates transcription of a nucleic acid sequence[, or gene,] coding for a polypeptide [to produce], wherein said[the] promoter or said[the] isolated promoter nucleic acid sequence comprises [corresponds to] a chimeric expression promoter or a promoter nucleic acid sequence according to claim 1, 6, or 49. [any one of claims 1 to 18 or 23 or 24.]

37) (Amended) The v[V]ector of[according to] claim 36, wherein said vector[it] is selected from the group consisting of the binary vectors pMRT1152, pMRT1171, pMRT1172, pMRT1185, pMRT1186, pMRT1187, pMRT1188, pMRT1182, pMRT1245, pMRT1246, pMRT1247, pMRT1248, pMRT1249, pMRT1250, pMRT1251, pMRT1252, pMRT1253 and pMRT1254.

38) (Amended) A t[T]ransgenic plant comprising [having] stably integrated into its genome at least one promoter or at least one promoter nucleic acid sequence according to claim 1, 6, or 49, [any one of claims 1 to 18 or 23 or 24 respectively.]

39) (Amended) The t[T]ransgenic plant of [according to] claim 38, wherein said plant[it] is one selected from dicotyledonous species comprising[, preferably] potato, tobacco, cotton, lettuce, tomato, melon, cucumber, pea, rape, beetroot, and[or] sunflower, or from monocotyledonous species comprising[, preferably] wheat, barley, oat, rice, and[or] corn.

40) (Amended) A p[P]ropagule of a transgenic plant according to claim [any one of claims] 38 or 39.

41) (Amended) The t[T]ransgenic plant propagule of[according to] claim 40, wherein it is a seed.

42) (Amended) A c[C]ell containing a promoter or a promoter nucleic acid sequence of claim 1, 6, or 49 [according to any one of claims 1 to 18 or 23 or 24 respectively], wherein said cell is selected from the group consisting of[and is preferably] a plant cell, human cell, animal cell, insect cell, bacterial cell, yeast cell, fungal cell, algal cell, and microalgal cell.

43) (Amended) The c[C]ell of[according to] claim 42, wherein it is a plant cell.

44) (Amended) A m[M]ethod for expressing a nucleic acid sequence[, or gene,] coding for a polypeptide [to produce,] by a cell, wherein said[the] method comprises[the steps consisting of]:

- transforming said[the] cell with a vector comprising at least one promoter or at least one promoter nucleic acid sequence [according to any one of claims 1 to 18 or 23 or 24] of claim 1, 6, or 49;

- culturing said [the] cell and expressing said polypeptide encoded by said sequence in said cell [under conditions enabling the expression of the nucleic acid sequence, or gene, coding for the polypeptide].

45) (Amended) The m[M]ethod of[according to] claim 44, wherein said[the] cell is a prokaryotic or an eukaryotic cell.

46) (Amended) The m[M]ethod of according to any one of claims claim 44 or 45, wherein said[the] cell is a cell selected from the group consisting of bacterial cells, fungal cells, yeast cells, insect cells, human cells, animal cells, algal cells, microalgal cells and plant cells.

47) (Amended) The m[M]ethod of according to any one of claims 44 to 46] claim 46, wherein said[the] cell is a plant cell.

48) (Amended) A m[M]ethod for [the] manufactur[e]ing[of] a transgenic plant of claim 38[according to any one of claims 38 or 39], or [of] a propagule of according to] claim 40, wherein said[the] method comprises[the steps consisting of]:

- transforming a plant cell with a vector comprising at least one promoter or at least one promoter nucleic acid sequence of according to any one of claims 1 to 18 or 23 or 24] claim 1, 6, or 49;

- selecting said[the] plant cell comprising[having] integrated said[the] promoter or said[the] promoter nucleic acid sequence;

- propagating said[the transformed] selected plant cell[, either] by culture or by regeneration of whole chimeric or transgenic plants.

49) (NEW) An isolated promoter nucleic acid sequence, comprising a fusion of a first sequence and a second sequence, wherein said first sequence and said second sequence comprise at least a part of each of the sequences with SEQ ID No. 1 and SE ID No. 2, respectively.

50) (NEW) The isolated promoter nucleic acid sequence of claim 23, wherein said first or second sequence comprises a sequence selected from the group consisting of sequences with SEQ ID Nos. 3-7 and 19-25.

51) (NEW) A directional deoxynucleotide building block for the construction of a chimeric expression promoter or an isolated promoter nucleic acid sequence according to claim 1, 6, or 23, wherein said sequence is selected from the group consisting of SEQ ID Nos. 8-14.